

IFW16

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/009,705E

DATE: 07/23/2004 TIME: 13:20:44

Input Set : A:\pto.lm.txt

Output Set: N:\CRF4\07232004\J009705E.raw

```
3 <110> APPLICANT: Ruprecht-Karls-Universitat Heidelberg
             Ruprecht-Karls-Universitat Heidelberg
      6 <120> TITLE OF INVENTION: Method for specifically detecting and identifying retroviral
nucleic
             acids/retro viruses in a specimen
W--> 8 <130> FILE REFERENCE: DE19921419.0
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- - 10 <140> CURRENT APPLICATION NUMBER: US/10/009,705E
    - 11 <141> CURRENT FILING DATE: 2001-11-28
    - 13 <160> NUMBER OF SEQ ID NOS: 70
    - 15 <170> SOFTWARE: PatentIn version 3.1
    - 17 <210> SEO ID NO: 1
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    - 20 <213> ORGANISM: Homo sapiens
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    - 23 <221> NAME/KEY: misc feature
    - 24 <222> LOCATION: (6)..(6)
    - 25 <223> OTHER INFORMATION: A or C or G or T
    - 28 <400> SEQUENCE: 1

### W--> 29 aragtnytdy chcmrggh 18

- 32 <210> SEQ ID NO: 2
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- 34 <212> TYPE: DNA
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- 39 <222> LOCATION: (1)..(1)
- 40 <223> OTHER INFORMATION: A or C or G or T
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- 47 <210> SEQ ID NO: 3
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- 50 <213> ORGANISM: Homo sapiens
- 52 <400> SEQUENCE: 3
- 53 tkkammskvy trcyhcargg g 21
- 56 <210> SEQ ID NO: 4
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- 58 <212> TYPE: DNA
- 59 <213> ORGANISM: Homo sapiens
- 61 <220> FEATURE:
- W--> 62 <221> NAME/KEY: SEQID5
  - 63 <222> LOCATION: (1)..(10)
  - 64 <223> OTHER INFORMATION:

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Input Set : A:\pto.lm.txt

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- 66 <220> FEATURE:
- 67 <221> NAME/KEY: misc feature
- 68 <222> LOCATION: (1)..(10)
- 69 <223> OTHER INFORMATION:
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  - 76 <211> LENGTH: 9
  - 77 <212> TYPE: DNA
  - 78 <213> ORGANISM: artificial sequence
  - 80 <220> FEATURE:
  - 81 <223> OTHER INFORMATION: arbitrary head sequence
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  - 84 <221> NAME/KEY: misc feature
  - 85 <222> LOCATION: (1)..(9)
  - 86 <223> OTHER INFORMATION: arbitrary head sequence
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  - 90 gaaggatcc 9
  - 93 <210> SEQ ID NO: 6
  - 94 <211> LENGTH: 90
  - 95 <212> TYPE: DNA
  - 96 <213> ORGANISM: Homo sapiens
  - 98 <220> FEATURE:
  - 99 <221> NAME/KEY: misc feature
  - 100 <222> LOCATION: (1)..(90)
  - 101 <223> OTHER INFORMATION: Capture probe 1A
  - 104 <400> SEQUENCE: 6
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  - 113 <213> ORGANISM: Homo sapiens
  - 115 <220> FEATURE:
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  - 117 <222> LOCATION: (1)..(90)
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  - 129 <212> TYPE: DNA
  - 130 <213> ORGANISM: Homo sapiens
  - 132 <220> FEATURE:
  - 133 <221> NAME/KEY: misc feature
  - 134 <222> LOCATION: (1)..(90)
  - 135 <223> OTHER INFORMATION: Capture probe 1E
  - 138 <400> SEQUENCE: 8

#### RAW SEQUENCE LISTING

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Input Set : A:\pto.lm.txt

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- 139 tatatcaact ctccggcttt gtgtcataat cttattcaga gtgatcttga tcacttttca 60
- 141 ctgccacaag atatcacact ggtccattad 90
- 144 <210> SEQ ID NO: 9
- 145 <211> LENGTH: 90
- 146 <212> TYPE: DNA
- 147 <213 > ORGANISM: Homo sapiens
- 149 <220> FEATURE:
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- 151 <222> LOCATION: (1)..(90)
- 152 <223> OTHER INFORMATION: Capture probe 1F
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- 158 acttctgcaa gatatcatgc tggtccatta 90
- 161 <210> SEQ ID NO: 10
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- 166 <220> FEATURE:
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- 168 <222> LOCATION: (1)..(90)
- 169 <223 > OTHER INFORMATION: Capture probe 1G
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- 175 cttccgcaag atatcacact gggtttgtta 90
- 178 <210> SEQ ID NO: 11
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- 180 <212> TYPE: DNA
- 181 <213> ORGANISM: Homo sapiens
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- 184 <221> NAME/KEY: misc feature
- 185 <222> LOCATION: (1)..(90)
- 186 <223> OTHER INFORMATION: Capture probe 1H
- 189 <400> SEQUENCE: 11
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- 192 cttccacaag atatcacact ggtccactac 90
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- 196 <211> LENGTH: 90
- 197 <212> TYPE: DNA
- 198 <213> ORGANISM: Homo sapiens
- 200 <220> FEATURE:
- 201 <221> NAME/KEY: misc\_feature
- 202 <222> LOCATION: (1)..(90)
- 203 <223> OTHER INFORMATION: Capture probe 1I
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- 209 ttccacaaga tatcacactg attgcctaca 90
- 212 <210> SEQ ID NO: 13
- 213 <211> LENGTH: 90
- 214 <212> TYPE: DNA

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DATE: 07/23/2004 TIME: 13:20:44

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Input Set : A:\pto.lm.txt

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286 <221> NAME/KEY: misc\_feature 287 <222> LOCATION: (1)..(90)

288 <223> OTHER INFORMATION: Capture probe 2E

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Input Set : A:\pto.lm.txt

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291 <400> SEQUENCE: 17 292 tttaaaaact cccctaccct ttttggggaa gccctccaac aggatcttat accattctga 60 294 gccagtaacc ctcactgcac tcttctccag 90 297 <210> SEQ ID NO: 18 298 <211> LENGTH: 90 299 <212> TYPE: DNA 300 <213> ORGANISM: Homo sapiens 302 <220> FEATURE: 303 <221> NAME/KEY: misc feature 304 <222> LOCATION: (1)..(90) 305 <223> OTHER INFORMATION: Capture probe 2F 308 <400> SEQUENCE: 18 309 tttaaaaatt ccgccaccct ttttggggaa gccctccaac aagatcttct accattctga 60 311 gccagtcccc ttaactgtaa ctcttcttca 90 314 <210> SEQ ID NO: 19 315 <211> LENGTH: 88 316 <212> TYPE: DNA 317 <213> ORGANISM: Homo sapiens 319 <220> FEATURE: 320 <221> NAME/KEY: misc feature 321 <222> LOCATION: (1)..(88) 322 <223> OTHER INFORMATION: Capture probe 2F 325 <220> FEATURE: 326 <221> NAME/KEY: misc feature 327 <222> LOCATION: (1)..(88) 328 <223> OTHER INFORMATION: Capture probe 2H 331 <400> SEQUENCE: 19 332 ttcaagaact cccccaccat ctttggggag gegttggctc gagacctcca gaagtttccc 60 334 accagagace taggetgegt gttgetee 88 337 <210> SEQ ID NO: 20 338 <211> LENGTH: 90 339 <212> TYPE: DNA 340 <213> ORGANISM: Homo sapiens 342 <220> FEATURE: 343 <221> NAME/KEY: misc feature 344 <222> LOCATION: (1)..(90) 345 <223> OTHER INFORMATION: Capture probe 2I 348 <400> SEQUENCE: 20 349 ttcaagaact cccttactat cttcggggag gctctgactt gagacttgca aaagtttcct 60 351 gctaaagacc taggctatgt cttgctcctg 90 354 <210> SEQ ID NO: 21 355 <211> LENGTH: 90 356 <212> TYPE: DNA 357 <213> ORGANISM: Homo sapiens 359 <220> FEATURE: 360 <221> NAME/KEY: misc\_feature 361 <222> LOCATION: (1)..(90)

365 <400> SEQUENCE: 21

362 <223> OTHER INFORMATION: Capture probe 2J

RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/10/009,705E

DATE: 07/23/2004 TIME: 13:20:45

Input Set : A:\pto.lm.txt

Output Set: N:\CRF4\07232004\J009705E.raw

#### Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; N Pos. 6

Seq#:2; N Pos. 1 L

#### VERIFICATION SUMMARY

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Input Set : A:\pto.lm.txt

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L:8 M:283 W: Missing Blank Line separator, <130> field identifier L:29 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0 L:44 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:0

L:62 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:4

L:71 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:4,Line#:64 L:71 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:4,Line#:69